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ORIGINAL PAPER

Effects of spatial pattern and relatedness in an experimental plant community

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Abstract Many plant species show limited dispersal resulting in spatial and genetic substructures within populations. Consequently, neighbours are often related between each other, resulting in sibling competition. Using seed families of the annuals *Capsella bursa-pastoris* and *Stachys annua* we investigated effects of spatial pattern (i.e. random versus aggregated) on total and individual performance at the level of species and seed families under field conditions. At the level of species, we expected that inferior competitors increase, while superior competitors decrease their performance within neighbourhoods of conspecifics. Thus, we expected a species by spatial pattern interaction. Sibling competition, however, might reduce the performance of competitors, when genetically related, rather than non-related individuals are competing. Therefore, aggregations at the level of seed families could decrease the performance of competitors. Alternatively, if the opposite outcome would be observed, kin selection might be hypothesized to have occurred in the past. Because heavy seeds are expected to disperse less than light seeds, we further hypothesized that kin selection might be more likely to occur in superior competitors with heavy, locally dispersed seeds (e.g. *Stachys*) compared to inferior competitors with light, more distantly dispersed seeds (e.g. *Capsella*). We found a significant species by spatial pattern interaction. Indeed, the inferior competitor, *Capsella*, showed increased reproductive biomass production in aggregated compared to random patterns. Whereas, the performance of the superior competitor, *Stachys*, was to some extent decreased by intra-specific aggregation. Although statistically not significant, effects of intrafamily aggregations tended to be rather negative in *Capsella* but positive in *Stachys*. Our results confirmed that spatial patterns affect growth and reproduction of plant species promoting coexistence in plant communities. Although, we could not provide strong evidence for sibling competition or kin selection, our results suggested that competition among relatives was more severe for *Capsella* (lighter seeds) compared to *Stachys* (heavier seeds).

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Introduction

Elucidating the many ways in which competing plant species manage to coexist is a major unresolved question in community ecology (Hutchinson 1961; Silvertown and Charlesworth 2001) and many hypotheses have attempted to explain the coexistence of species with similar life histories (see, e.g. Shmida and Ellner 1984; Chesson 2000a, b; Wright 2002; Barot 2004; Silvertown 2004). However, explaining how large numbers of competing plant species manage to coexist in the absence of obvious niche differentiation, conspicuous life history trade-offs (e.g. competition/colonization trade-off; Rees 1995; Turnbull et al. 1999, 2004) or evident disturbance (Connell 1978) remains challenging and controversial (see, e.g. Amarasekare 2003; Barot 2004; Silvertown 2004).

As a consequence of limited seed dispersal and/or clonal growth, most plant species aggregate intraspecifically, thereby increasing the importance of intra- versus interspecific competition (spatial segregation hypothesis, Pacala 1997). This is particularly relevant for sessile organisms where survivorship and fecundity are most affected by local population density rather than by the average global population density (Pacala and Silander 1985; Pacala 1997; Murrell et al. 2001). There is a large body of theory that underlines the importance of spatial pattern for ecological phenomena, such as coexistence and maintenance of biodiversity (Kareiva 1990; Bergelson 1990; Coomes et al. 2002; Bolker et al. 2003; Levine and Murrell 2003). However, compared to the large body of theory, there is still surprisingly little empirical evidence for the importance of spatial structure in shaping plant communities. Nevertheless, some experiments have shown that intraspecific aggregation might foster coexistence by allowing inferior competitors to increase their fitness (e.g. seed production). This might increase their persistence in the plant community and slow down competitive exclusion (Schmidt 1981; Bergelson 1990; Stoll and Prati 2001; Monzeglio and Stoll 2005). For example, if disturbances create gaps, inferior competitors might be able to colonize these gaps as long as they can produce enough seeds somewhere in the community. The few experimental studies that are available manipulated the spatial arrangement in an agricultural context (e.g. Harper et al. 1961; Brophy and Mundt 1991; Stauber et al. 1991; Norris et al. 2001). However, to our knowledge only one focused on natural communities (Turnbull et al. 2007). Therefore, the question to what extent and under which conditions intraspecific aggregation of species promotes coexistence by slowing down competitive exclusion remains controversial (Chesson 1991; Chesson and Neuhauser 2002; Murrell et al. 2002; Bolker et al. 2003).

Spatially limited seed dispersal can lead to pronounced aggregation of conspecifics (see, e.g. Seidler and Plotkin 2006), with the additional consequence that neighbours are often genetically related (half- or full-siblings) to each other. In other words, when the prevailing pattern of dispersal results in relatives being aggregated in space and interacting primarily with one another, then local competition may become sibling competition (Cheplick 1992, 1993a, b; Kelly 1996; Lambin et al. 2001). Therefore, in plant species where dispersal is limited, the effects of spatial patterns (i.e. intraspecific aggregation) should not only operate at the level of species but also at the level of genotypes within species (Schmid 1990; Vuorisalo et al. 1997; Wilson and Nisbet 1997).

In general, the intensity of competition is thought to increase with genetic similarity of the competitors (Maynard Smith 1978). This hypothesis, known as the resource-partitioning hypothesis (Young 1981; Argyres and Schmitt 1992), states that genetically variable offspring will experience less severe competition than genetically similar or identical offspring, because more diverse offspring are predicted to show greater ability to partition limiting resources. In addition, parents will leave fewer offspring if there is severe competition between their descendants (Maynard Smith 1978). Therefore, sibling competition can be broadly considered as negative interactions between genetically related individuals and should in principle be avoided (Cheplick 1992, 1993a, b).

Conversely, relatedness of individuals in a population may lead to kin selection (Hamilton 1964). The kin selection hypothesis predicts that individuals will behave altruistic, when $rb - c > 0$, where c is the fitness cost to the altruist, b the fitness benefit to the beneficiary and r is their genetic relatedness. In other words, an individual may behave altruistic (e.g. less competitive towards their relatives), even if the altruistic behaviour reduces its own fitness, if the costs are compensated with increased fitness of its relatives. In plant populations, kin selection may be a significant evolutionary force that counteracts sibling competition, because many species have limited seed dispersal (Goodnight 1985; Goodnight and Stevens 1997). Especially in plant species with heavy, locally dispersed seeds, local aggregation will increase sibling competition, but at the same time increase the potential for kin selection. On the other hand, in plant species with light far-dispersed seeds, seedlings are expected to be less aggregated and therefore sibling competition might occur less frequently. Thus, in plant species with frequent sibling interactions the possibility that kin selection might counteract potentially negative effects should not be excluded (Nakamura 1980; Cheplick 1993b; Kelly 1996; Griffin and West 2002). However, despite the straightforward predictions of the sibling competition hypotheses, few empirical tests have been made and the results are inconsistent. Amongst the studies, which investigated sibling competition in plants (Cheplick 1992), some showed that plants competing with genetically related individuals perform worse than plants competing with unrelated individuals (Willson et al. 1987; Kelley 1989; Argyres and Schmitt 1992; Karron and Marshall 1993). Others provide evidence that plants competing with genetic relatives outperform those competing with unrelated individuals (Willson et al. 1987; Tonsor 1989; Andalo et al. 2001; Donohue 2003). Hence, so far it remains difficult to unequivocally argue which of the both hypotheses is more likely to describe the dynamics of local competition in plant populations and communities.

The experiment reported here extends a previous spatial pattern experiment (Monzeglio and Stoll 2005) that assessed effects of intraspecific aggregation on species performance in experimental plant communities. The present experiment went further and aimed to evaluate effects of intraspecific aggregation at the level of seed families within species. Specifically, we collected seed families, which allowed us to assess not only effects of intraspecific aggregation but also effects of intrafamily aggregation on the growth and fitness of individuals. Based on our previous experiments, we expected that weak or competitively inferior species (e.g. *Capsella bursa-pastoris*) would produce more biomass when locally aggregated compared to randomly distributed and that intraspecific aggregation decreased biomass production of strong or competitively superior species (e.g. *Stachys annua*). In statistical terms, we expected a species by spatial pattern interaction.

Based on the sibling competition and the associated resource-partitioning hypothesis, we further expected that groups of relatives (i.e. intrafamily aggregation) would generally produce less vegetative and reproductive biomass compared to groups of non-relatives (i.e. intraspecific aggregation). In other words, biomass production in the intrafamily

aggregation would be lower compared to the intraspecific aggregation. In contrast, if the opposite pattern would occur, then the operation of kin selection might be indicated. In this case, the performance of genetically related individuals would increase from intraspecific to intrafamily aggregations. Because *Capsella* has small and lighter seeds compared to *Stachys*, the former is expected to disperse its seeds less locally than the latter. Therefore, we hypothesized that kin selection might be more likely in *Stachys* where sibling competition should be more frequent compared to *Capsella*. Consequently we expected that intrafamily aggregation might be more likely to have positive effects in *Stachys* than *Capsella*.

Materials and methods

The purpose of this experiment was to investigate effects of spatial patterns (random versus aggregated) at the level of species (intraspecific aggregation) and seed families (intrafamily aggregation) within a species mixture under experimental field conditions [at the Research Institute of Organic Agriculture (FiBL), Frick, Switzerland]. We used four annual plant species (*C. bursa-pastoris*, *S. annua*, *Stellaria media* and *Poa annua*) representing common species composition in natural communities. *Capsella* and *Stachys* were the target species, while *Poa* and *Stellaria* were the matrix species. Seeds of the target species were collected from single mother plants grown in a previous experiment (Monzeglio and Stoll 2005) with a similar design as the present one. The mother plants were grown under four maternal treatments with different combinations of two spatial pattern (random versus aggregated) and two densities (low versus high). Seeds of the matrix species were obtained from a commercial seed supplier (Herbiseed, Wokingham, Berkshire, UK).

Plant species

Capsella bursa-pastoris L. (Brassicaceae) is a widely distributed, cosmopolitan annual pioneer species of disturbed ground and trampled sites. It forms rosettes with a multi-flowered erect stem up to 40 cm high. Flowers are usually self-pollinated; however small insects (e.g. flies, small bees) visit the flowers. In disturbed European sites, greater levels of genetic heterogeneity have been recorded for populations sampled from recently cultivated than from non-cultivated sites (Bosbach and Hurka 1981), suggesting a positive relationship between the degree of environmental variability and genetic variation. *Capsella* reproduces entirely by seeds, which are small and light, dispersed by wind or rain drops. The majority of the seeds usually fall between 15 and 30 (–50) cm of the parent plant (Aksoy et al. 1998). *Stachys annua* L. (Lamiaceae) is an annual species with a multi-flowered erect stem up to 40 cm high. This species grows for example in corn and tubercrop fields and gravel-pits. It prefers warm, dry and calcareous soils. *Stachys* is quite rare in most parts of Europe. It is predominately outcrossed and flowers from June to October. The hermaphroditic flowers are usually pollinated by hymenoptera, lepidoptera, or diptera. Although birds may disperse some seeds, *Stachys* seeds have no special structures aiding dispersal (e.g. elaisomes) and most of the relatively heavy seeds probably fall down in very close proximity of their mother plants. *Stellaria media* L. (Caryophyllaceae) is prostrate to ascending, with high-adventitious rooting and a height of up to 40 cm. *Poa annua* L. (Poaceae) has adventitious roots at the first nodes and tillers up to 30 cm high and reproduces primarily by seeds. *Stellaria* and *Poa* are widely distributed,

cosmopolitan annuals of disturbed habitats. Generally, *C. bursa-pastoris*, *P. annua* and *S. media* occur together.

Collection of seed families

Seed families of the target species *Capsella* and *Stachys* used in the present experiment were collected from mother plants grown in a previous spatial pattern experiment (see Monzeglio and Stoll 2005 for a detailed description of the experimental design) in which all four species (*Capsella*, *Stachys*, *Poa* and *Stellaria*) were grown in combinations of two different spatial patterns (random versus aggregated) and two different densities (low versus high). Since *P. annua* and *S. media* did not produce enough seeds, for the present experiment seeds were obtained from a commercial supplier (Herbiseed). Seeds of *Capsella* and *Stachys* were collected in summer 2002 and the maternal environments (spatial pattern and density) recorded. All seeds from a single mother plant are referred to as seed family and are at least half-sibs. We started the seed collection of *Capsella* on August 8, and mature seeds were collected daily until August 15. The last collection was done on September 11 when collection of the *Stachys* seeds started. In this case, in order to avoid too much seed loss, we collected the whole plant and separated the seeds later in the laboratory. These collections were done on September 20, October 1 and 8 and November 12. Seeds of each mother plant were counted mechanically (seed counting machine, Elomor) and the seed families with the largest number of seeds were chosen for the present experiment. In total, 36 seed families of *Capsella* (nine from random and aggregated low densities, and 18 from random high density) were used. For *Capsella* in aggregated high density no seed families produced enough seeds for the present experiment. As a result for the high density we had only mother plants grown in random spatial patterns. For *Stachys*, 36 seed families (nine from each of the four maternal treatment combinations) yielded enough seeds for the present experiment. Because the experiment had to be started, only the remaining seeds could be counted and weighed. The mean seed mass of *Capsella* was much lower than that of *Stachys* (Table 1). *Capsella* grown in aggregated patterns produced heavier seeds than *Capsella* grown in random spatial patterns ($t = 2.21$, $df = 25$ and $P = 0.037$). *Stachys* did not show any significant differences in seed weight between mothers grown in random versus aggregated spatial patterns or at high versus low density. There was no significant interaction between maternal density and maternal spatial pattern ($F_{1,35} = 1.4$; $P = 0.253$).

Experimental design

The experiment was designed as randomized split-plot and contained two blocks (0.6×6.0 m, separated by 0.5 m) each with six main plots (0.6×0.6 m, separated by 0.3 m). The three spatial pattern treatments (see below) were used as plot-level treatments yielding four replicates per treatment (two per block). The plots were subdivided into nine subplots (0.2×0.2 m) that were divided in an upper and lower half, in such a way that each half contained 4 1/2 subplots (the central subplot was halved) (Fig. 1A).

In order to avoid any misunderstanding about the spatial patterns of the present experiment (i.e. random, intraspecific and intrafamily aggregation) and spatial patterns in

Table 1 Mean seed mass (mg) of *Capsella bursa-pastoris* and *Stachys annua* across seed families ($N = 9$) collected from individual mother plants grown in two different spatial patterns (random versus aggregated) and at two densities (low versus high)

Maternal density	Maternal spatial pattern	<i>Capsella bursa-pastoris</i>			<i>Stachys annua</i>		
		<i>N</i>	Seed mass	<i>SE</i>	<i>N</i>	Seed mass	<i>SE</i>
Low	Random	9	0.109	0.002	9	2.84	0.26
	Aggregated	9	0.117	0.004	9	2.75	0.15
	Mean		0.113	0.002		2.79	0.15
High	Random	9	0.107	0.004	9	3.05	0.31
	Aggregated	0			9	2.37	0.28
	Mean		0.107	0.004		2.71	0.22
Mean	Random	18	0.108	0.002	18	2.94	0.20
	Aggregated	9	0.117	0.004	18	2.56	0.16

For each spatial pattern/density combination nine different seed families were used (except for *Capsella* which had no seed families that produced enough seeds in high densities and aggregated pattern). For *Capsella* in high density and random pattern only nine, instead of 18, seed families had enough remaining seeds to be counted and weighed

N number of mother plants or seed families, *SE* standard error

which the mother plants grew (i.e. random and intraspecific aggregation, see Experimental design for maternal environment) we describe the main experimental design and the design corresponding to the maternal environment separately. We start with the spatial pattern of the present experiment.

In the present experiment, we used the four-species mixture and one density level (i.e. 100 seeds per cell, for a total of 40,000 seeds/m²). The three spatial pattern treatments were attained as follows. In the random pattern, 44 seeds of each of the nine seed families of *Capsella* and *Stachys* (i.e. for a total of ~400 seeds for each species) together with 400 seeds of *Poa* and 400 seeds of *Stellaria* were mixed and randomly scattered over the subplots, such that the individuals experienced inter- and intraspecific as well as intrafamily encounters at similar frequencies. In the aggregated patterns (i.e. intraspecific and intrafamily aggregated patterns), the subplots were further subdivided into 16 cells (5 × 5 cm), and each cell contained only one species in such a way that individuals experienced more intra- than interspecific encounters. For the two species with seed families (*Capsella* and *Stachys*) seeds were aggregated at two levels. The first level consisted of groups of conspecifics, and will be referred to as intraspecific aggregation. In the intraspecifically aggregated pattern each cell contained ~100 seeds made up of 11 seeds from each of the nine seed families. The second level was made up of individuals of the same seed families and will be referred to as intrafamily aggregation. In the intrafamily aggregated pattern each cell contained 100 seeds from the same seed family in such a way that individuals in the centre of the cells competed only with genetically related individuals (Fig. 1A). Two seed families were used per species and subplot. The species and seed families were randomly allocated to the cells. For the species without seed families (*Poa* and *Stellaria*) each cell always contained 100 seeds sown as the intraspecific aggregations.

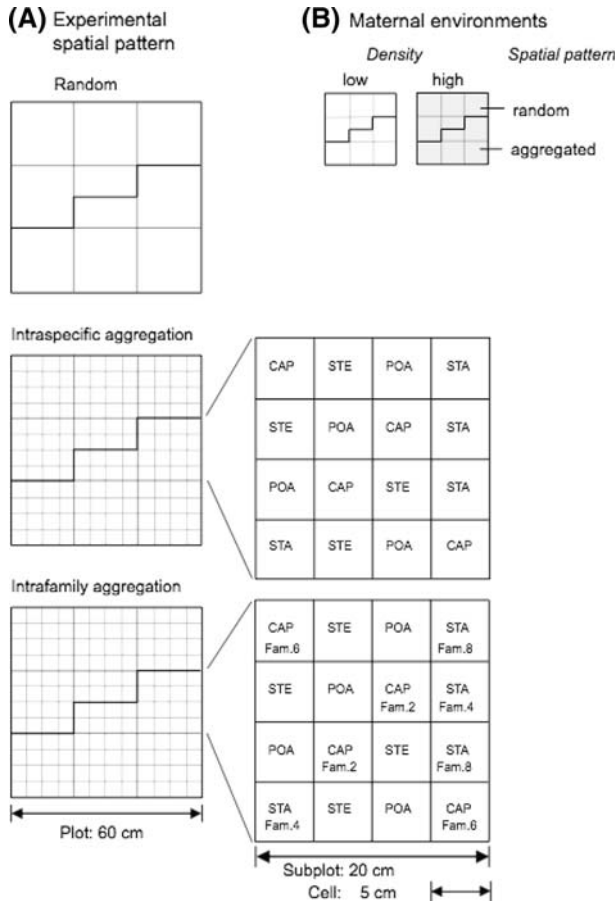


Fig. 1 Experimental design. **(A)** Experimental spatial pattern. Three different experimental spatial patterns, which varied at the plot (60 × 60 cm) level, were realized as follows. In the random spatial pattern, 44 seeds from each of nine families of *Capsella* (CAP) and *Stachys* (STA) and 400 seeds of *Poa* (POA) and *Stellaria* (STE) were mixed and scattered over the subplots (20 × 20 cm). In the intraspecific aggregation, subplots were further divided into 16 cells (5 × 5 cm). Four randomly selected cells per species were sown with 11 seeds from each of nine families of *Capsella* or *Stachys* or 100 seeds of *Poa* or *Stellaria*. In the intrafamily aggregated pattern, two randomly selected cells were sown with 100 seeds from one seed family and two with 100 seeds from another seed family of *Capsella* or *Stachys*. The other two species (*Poa* and *Stellaria*) were sown as in the intraspecific aggregations. Spatial pattern were replicated four times (two randomly assigned within each of two blocks). **(B)** Maternal environments. The mother plants, from which seed families of *Capsella* and *Stachys* were collected, grew either in low or high density and random or aggregated patterns of a previous experiment. The maternal density (i.e. low versus high) was varied at the plot level, while the maternal spatial pattern (random versus aggregated) at the 'half-plot' level. One half (4 1/2 subplots) of each plot was sown with seeds from mothers grown in a random spatial pattern and the other half was sown with seeds from mothers grown in an aggregated spatial pattern. The maternal spatial pattern was randomly assigned

Experimental design for maternal environment

The density in which the mother plants grew was used as plot-level treatments. Each block contained six plots, i.e. three plots with seeds from mother plants grown at low and three

plots with seeds from mother plants grown at high density. The spatial pattern in which the mothers grew was used as within-plot treatment (i.e. half-plot). Randomly allocated lower or upper halves were either sown with seeds from mothers grown in random or spatially aggregated patterns (for *Capsella* this was only possible for seeds from low-maternal densities and all seeds from mothers grown in high densities were from mothers grown in random spatial patterns) (Fig. 1B).

Additional experimental settings

In order to assess the germination activity of the seed families, simple germination trials with and without Gibberellic acid (0.01%) both in the laboratory and in the field were carried out in May 2003. These experiments showed low-germination rates of field collected seeds without Gibberellic acid. Therefore, to increase germination, seeds of *Capsella* and *Stachys* were soaked in a solution of Gibberellic acid at 4–5°C for three to 5 days before sowing.

The blocks were established on May 16 and harvested in fall 2003. During the first month the two blocks were covered with a plastic tunnel (200 holes/m², GVZ-Bolltec, AG, Zurich, Switzerland) to protect the seedlings against sunlight and adverse weather. The tunnel was never completely closed, 20–30 cm from the ground were left open to permit the air to circulate and in July the tunnel was removed completely. Because of the heavy soil (high-clay content), each plot was covered with a layer (2 cm) of commercial garden soil before sowing to increase seed germination. The plots were sown between June 16 and 19 and watered with an automatic irrigation system (Gardena AG, Bachenbülach, Switzerland). The system was programmed to give rain-cycles of 1 min (~1 l water) starting at 5.15, 6.15, 7.15 a.m., and 7.15, 8.15, 9.15 p.m. The duration of the 9.15 p.m. rain-cycle was changed from 1 to 2 min on June 26. Because of the extremely hot summer 2003 on July 2, one additional nozzle was positioned in the middle of each plot. These additional nozzles were programmed to give rain-cycles lasting 2 min starting at 8.30, 9.30, 12.30 a.m., and 4.30, 8.30, 9.30 p.m. and removed on July 29. After sowing, the seeds were covered by sieving a thin layer of commercial garden soil, which was slightly pressed down to prevent the seeds from being washed away. A snail fence enclosed the entire experimental field.

Harvest

The aboveground biomass of the four species was harvested at the subplot level (0.2 × 0.2 m) between September 22 and October 15. Additionally, for the two species with seed families (*Capsella* and *Stachys*) the harvested biomass was separated into vegetative and reproductive parts and the total number of individuals was counted. Because *Capsella* started to produce mature seeds earlier, mature seeds were collected regularly from 25 July to 15 August 2003, and then added to the final reproductive biomass. The harvested biomass was dried for 12 h at 60°C and then stored. Before weighing, the biomass was dried again for 4–5 h at 60°C.

Statistical analysis

The data from *Capsella* and *Stachys* were analysed with a multi-factorial analysis of variance (ANOVA) for a split-plot design. Since maternal density, which varied at the plot level, never showed significant effects it was excluded from the analysis and ANOVA tables. Similarly, because maternal spatial pattern, which varied at the half-plot level, never showed significant effects, we pooled the data from the subplots at the plot level. That is, we summed the biomass and number of individuals of the nine subplots, and used the plots ($N = 12$ with four replicates per spatial pattern) as experimental units. This pooling allowed us to simplify the ANOVA tables and the communication of the results but it did not change any of the reported results. Since we had no seed families for *Poa* and *Stellaria*, these species were considered as matrix species and not further analysed.

The spatial pattern treatments were tested against the plot-level residual mean square. In addition we used linear contrasts to separate them into (1) random versus aggregated (i.e. intraspecific and intrafamily aggregation), (2) intraspecific versus intrafamily aggregated pattern. There was one plot with a random spatial pattern treatment, in which the plants germinated particularly poorly or died soon after they germinated because of the extremely hot and dry weather. This was the only plot in which *Capsella* produced $<10 \text{ g/m}^2$ and the only plot in which biomass production of *Stachys* was less than the mean $\pm 3 \text{ SE}$. As it turned out that *Capsella* produced less biomass in plots with random spatial patterns than aggregated plots, including this plot in the *Capsella* analysis rendered all effects more significant than reported. On the other hand, *Stachys* produced most biomass in the plots with random spatial patterns, except this particular one. Including this plot in the *Stachys* analysis increased the residual variation so much that all effects for *Stachys* became non-significant. Because the results of the analysis, especially for *Stachys*, changed significantly depending on whether we considered this plot or set the plot values as missing, we decided to present both analyses. The analysis in which the values for this unusual random plot were omitted is presented in the results (Table 3) and used to plot the charts. The analysis using all plots is presented as appendix.

We counted the number of seedlings 1 month after the beginning of the experiment and tested it as covariate. The covariate, however, had no significant effect and was consequently omitted. Even though the maternal environments had no significant effect on the main spatial pattern effect, we also tested initial seed mass for *Stachys* and *Capsella* as covariate. Again, the covariate had no significant effect and was omitted.

All data were expressed as number of individuals or grams per square metre for total biomass production or grams per individual for average biomass production. All biomass data were logarithmically transformed [$\log_{10}(Y)$] to obtain normal distribution of the residuals and homogeneity of variances. Mean and SE from the analyses (back-transformed in the case of biomass) are presented throughout. All analyses were conducted using GENSTAT 5 (Payne et al. 1987).

Results

First of all, both species differed significantly in their overall performance (Table 2). Generally, total biomass production of *Capsella* was much lower ($18.9 \pm 4.0 \text{ g/m}^2$, average $\pm \text{SE}$ over 11 plots) than that of *Stachys* ($1,166.7 \pm 30.1 \text{ g/m}^2$). The matrix species *Stellaria* produced $54.2 \pm 10.7 \text{ g/m}^2$ while *Poa* did not germinate.

Table 2 Results of ANOVA combining both species *Capsella bursa-pastoris* and *Stachys annua* omitting one unusual random plot

Source of variation	df (m.v.)	Total		Average			
		Number of individuals		Vegetative biomass		Reproductive biomass	
		F	P	F	P	F	P
Block	1	6.4	0.039	0.0	0.964	1.0	0.353
Spatial pattern (SP)	2	3.3	0.100	1.8	0.236	7.3	0.019
Random versus aggregated ^a	1	4.8	0.064	3.0	0.129	14.4	0.007
Intraspecific versus intrafamily aggregated ^a	1	1.7	0.235	0.6	0.459	0.3	0.63
Plot level	7 (1)	0.6		1.3		1.1	
Species (S)	1	69.1	<0.001	322.9	<0.001	480.2	<0.001
S × SP	2	2.9	0.112	3.9	0.067	9.5	0.008
S × random versus aggregated ^a	1	5.3	0.051	6.5	0.034	18.6	0.003
S × intraspecific versus intrafamily aggregated ^a	1	0.6	0.478	1.2	0.301	0.5	0.512
Residual	8 (1)						
Total	21 (2)						

Effects of spatial patterns (random versus two levels of aggregation) on total (sum over plots) number of individuals, total and average (per individual) aboveground vegetative and reproductive biomass production were tested against plot-level residuals

df degrees of freedom, (m.v.) missing values, F F-value (variance ratio), P error probability

^a Linear contrasts to separate the effects of spatial pattern into differences between random versus aggregated (i.e. intraspecific and intrafamily aggregation) and intraspecific versus intrafamily aggregation effects

Table 3 Results of ANOVA for *Capsella bursa-pastoris* (top) and *Stachys annua* (bottom) omitting one unusual random plot

Source of variation	df	(m.v.)	Total	Average									
				Number of individuals		Vegetative biomass		Reproductive biomass		Vegetative biomass		Reproductive biomass	
				F	P	F	P	F	P	F	P	F	P
<i>Capsella bursa-pastoris</i>													
Block	1	1.1	0.327	0.0	0.982	0.8	0.415	6.7	0.036	17.9	0.004		
Spatial pattern	2	1.4	0.303	2.5	0.150	7.9	0.016	11.1	0.007	33.9	<0.001		
Random versus aggregated ^a	1	1.3	0.288	4.2	0.079	15.5	0.006	21.9	0.002	67.6	<0.001		
Intraspecific versus intrafamily aggregated ^a	1	1.5	0.257	0.8	0.394	0.3	0.576	0.3	0.631	0.2	0.683		
Residual	7	(1)											
Total	10	(1)											
<i>Stachys annua</i>													
Block	1	5.3	0.056	0.0	0.847	2.2	0.185	5.3	0.056	7.3	0.031		
Spatial pattern	2	3.2	0.103	6.1	0.030	1.9	0.218	1.3	0.329	1.8	0.234		
Random versus aggregated	1	5.2	0.056	10.8	0.013	2.9	0.133	1.1	0.321	2.0	0.199		
Intraspecific versus intrafamily aggregated	1	1.2	0.316	1.3	0.293	0.9	0.368	1.5	0.264	1.6	0.247		
Residual	7	(1)											
Total	10	(1)											

Effects of spatial patterns (random versus two levels of aggregation) on total (sum over plots) number of individuals, total and average (per individual) aboveground vegetative and reproductive biomass production were tested against plot-level residuals

df degrees of freedom, (m.v.) missing values, *F* *F*-value (variance ratio), *P* error probability

^a Linear contrasts to separate the effects of spatial pattern into differences between random versus aggregated (i.e. intraspecific and intrafamily aggregation) and intraspecific versus intrafamily aggregation effects

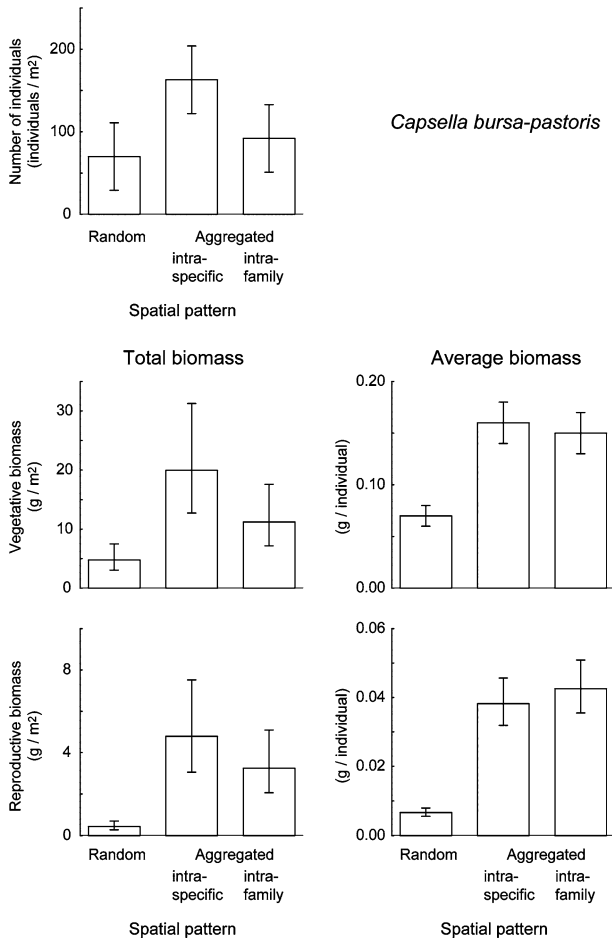
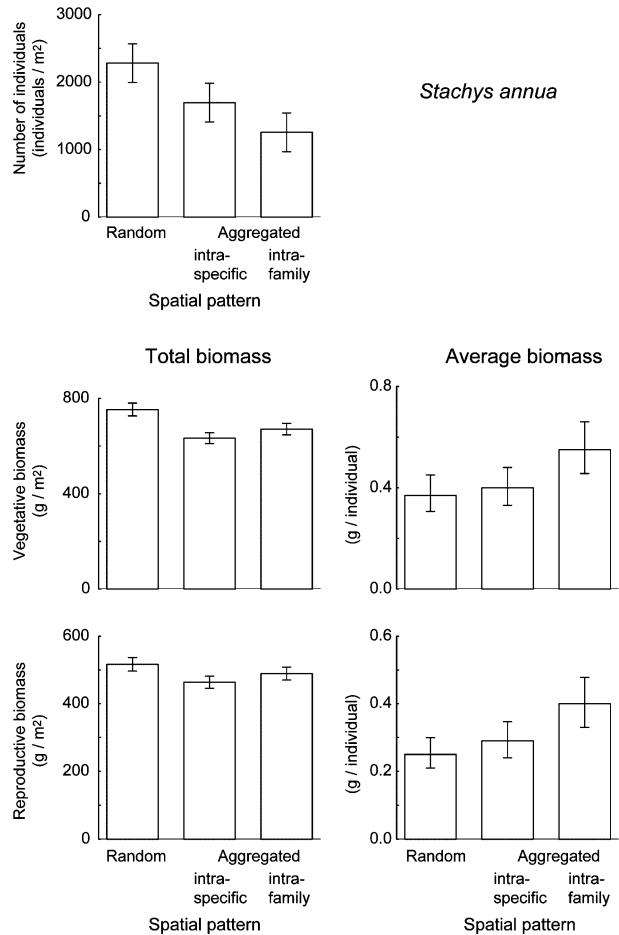


Fig. 2 Number of *Capsella bursa-pastoris* individuals (top), total (left, g/m²) and average (right, g per individual) aboveground vegetative (middle) and reproductive (bottom) biomass grown in random, intraspecific or intrafamily aggregated spatial patterns. The bars represent mean \pm 1 SE from ANOVA for number of individuals and backtransformed means from $\log_{10}(Y)$ -transformed data from ANOVA's for vegetative and reproductive biomass

The analysis including both species revealed a significant spatial pattern effect for total reproductive biomass and average vegetative and reproductive biomass (Table 2). For example, averaged over both species there was significantly more total reproductive biomass in the aggregated (intraspecific 46.9 g/m²; intrafamily 39.9 g/m²) than in the random patterns (15.2 g/m²). There was a marginally significant species by spatial pattern interaction for total and average vegetative biomass and a significant interaction for total and average reproductive biomass (Table 2). The linear contrast showed that the main differences were between the random compared to the aggregated patterns.

Because the spatial pattern influenced the performance of the two species differently, we present species-specific results separately and begin with *Capsella*. *Capsella* generally had fewer individuals and produced less total and average biomass in the random compared to the aggregated patterns. The main spatial pattern effect was significant for total

Fig. 3 Number of *Stachys annua* individuals (top), total (left, g/m²) and average (right, g per individual) aboveground vegetative (middle) and reproductive (bottom) biomass grown in random, intraspecific or intrafamily aggregated spatial patterns. The bars represent mean \pm 1 SE from ANOVA for number of individuals and backtransformed means from log₁₀(Y)-transformed data for vegetative and reproductive biomass



reproductive and average vegetative and reproductive biomass. The linear contrasts between random versus intraspecific and intrafamily aggregated pattern were highly significant. However, the differences between intraspecific versus intrafamily aggregated pattern were not significant (Table 3, Fig. 2). Although *Capsella* produced ~70% less total vegetative biomass in the random compared to both (pooled) aggregated pattern, this effects was only marginally significant. The number of individuals was not significantly affected by the spatial pattern (Table 3, Fig. 2).

Stachys was less affected by the spatial pattern than *Capsella*. Nevertheless, *Stachys* had more but on average smaller individuals in random compared to both aggregated patterns. The main spatial pattern effect was not significant for the number of individuals, but the linear contrasts showed that *Stachys* had slightly more individuals in the random compared to the intraspecific and intrafamily aggregated pattern (Table 3, Fig. 3). Total vegetative biomass production of *Stachys* was significantly affected by the spatial pattern, with more total vegetative biomass in the random compared to both intraspecific and intrafamily

aggregated pattern (Table 3, Fig. 3). The average vegetative biomass was not significantly affected by the spatial pattern. Total and average reproductive biomass of *Stachys* was not affected by the spatial pattern.

Discussion

Based on aboveground biomass production we defined *S. annua* as a strong and *C. bursa-pastoris* as a weak competitor. The weak competitor produced much more reproductive biomass per individual in aggregated compared to random spatial patterns. Because they also tended to be more numerous in aggregated compared to random spatial patterns, total reproductive biomass production was much higher in aggregated compared to random spatial patterns. Individuals of the strong competitor in contrast, were more numerous in random compared to the aggregated spatial patterns. However, individuals in random patterns tended to produce less vegetative and reproductive biomass per individual compared to individuals in aggregated patterns. Thus, *Stachys* almost exactly compensated for the different number of individuals such that there were only small and insignificant differences in the production of total reproductive biomass between the different spatial patterns (516 g/m² in random, 463 and 488 g/m² in intraspecific and intrafamily aggregated patterns, respectively). Interestingly, our findings showed, averaged over both species, an overall increase in total reproductive biomass in the aggregated compared to random pattern. This suggested that *Capsella* produced overproportionally more total reproductive biomass in the aggregated pattern than *Stachys* in the random pattern. This result further supports the notion that spatial structure seems to be relatively more important for small seeded (e.g. *Capsella*) compared to large seeded species (e.g. *Stachys*) (Monzeglio and Stoll 2005; Turnbull et al. 2007).

Our results for the weak competitor were generally consistent with previous findings (Stoll and Prati 2001; Monzeglio and Stoll 2005). They support our expectation that weak competitors increased their fitness (e.g. survival and seed production) within neighbourhoods of conspecifics compared to neighbourhoods of heterospecifics, at least in the short run. In addition, data on *Stachys* showed that intraspecific aggregation decreased the number of individuals and to some extent the performance of strong competitors. Thus, our results at the species level generally agreed with our earlier work on the effects of spatial patterns on species performance.

However, experimental studies like ours have the limitation that the absolute and relative densities of the species, the structure and scale of the plant community and the strength of competition are determined by the experiment rather than natural processes. In real communities, the degree of spatial aggregation is often less extreme and the spatial structure is more complex. Turnbull et al. (2007) quantified the effects of spatial structure on individuals, population and community biomass within a natural community of annuals. They found that the effects of spatial structure on total community biomass were relatively small. This was because common species have generally weak spatial structure and they draw down the effect on the community as a whole. Therefore, intraspecific aggregation as process that may delay competitive exclusion would be far more important for rare and weak (e.g. small seeded species) than for common and strong (e.g. large seeded species) competitors. Our results lend further support to this general conclusion, because spatial pattern effects for the small seeded species *Capsella* were much stronger than for the large seeded species *Stachys*.

Our results at the level of seed families, however, were less clear and not statistically significant. Nevertheless, our results suggested, that effects of intrafamily aggregation differed depending on the species. Groups of relatives in the small seeded species (*Capsella*) tended to be less productive compared to groups of non-relatives in producing biomass. In particular, total reproductive biomass production in intrafamily aggregated patterns was only about half of that in intraspecifically aggregated patterns. This agrees with predictions from the sibling competition hypothesis and suggests that groups of relatives in *Capsella* indeed have a lower potential for resource partitioning than groups of non-relatives. For the large seeded species (*Stachys*), groups of relatives were slightly more productive compared to groups of non-relatives. Although the differences were not statistically significant, the general pattern agreed with our hypothesis that kin selection might be more likely to occur in plant species with heavy, locally dispersed seeds. Cheplick and Kane (2004) did a similar greenhouse experiment using *Triplasis purpurea* growing either alone or in inter- or intrafamily competition (note that what they call interfamily competition corresponds to our intraspecific aggregated pattern). Their findings generally support the resource partitioning hypothesis. Similar to other studies (e.g. Taylor and Aarssen 1990; Donohue 2003), they showed, however, that the effects of neighbour relatedness on the growth and fitness differed among families. This suggests genetic variation of competitive ability among seed families. Contrary to Cheplick and Kane (2004), we could not obtain detailed information on the different competing seed families and our experiment was conducted under field conditions. To obtain information on the different competing seed families in the intrafamily aggregated pattern, we would have had to harvest the biomass at the cell level (5×5 cm) rather than the subplot level (20×20 cm). In addition, in the intraspecific and random pattern we would have had to label each individual seed in order to recover its family at harvest. This was impossible. However, we cannot exclude the possibility that one or more seed families performed better in the intrafamily compared to the intraspecific aggregation even though *Capsella*, on average, decreased its fitness in the intrafamily compared to the intraspecific aggregated pattern.

The very high density used in our experiment (40,000 seeds/m²) might be one reason for the prevailing effect of resource partitioning. Koelewijn (2004) investigated the effects of different densities on competing seed families and showed that density had stronger negative effects in absolute terms on the performance of the seed families and that the consequences of sibling competition depend on the frequency and relatedness of neighbours. Escarré et al. (1994) examined density effects and neighbour relatedness in a sib/non-sib competition experiment on the clonal *Rumex acetosella*. They found density-dependent effects of the degree of relatedness between plant individuals. At low densities, there were no growth differences, but when the density was doubled, the absolute sexual biomass was higher in non-sibs than in sib treatments, suggesting that competition was stronger between related plants. Therefore, we speculate that in our experiment, a lower density might have favoured the positive intrafamily effects observed in *Stachys*. In contrast, a higher density might have increased the negative effect of sibling competition observed in *Capsella*. Moreover, there is evidence that the effect of high relatedness may be modulated by the scale of competition (e.g. Queller 2004, Griffin et al. 2004, Frank 1998). Even if highly localized dispersal is likely to promote the evolution of altruistic behaviour (Hamilton 1964; Wilson 1987), the scale of competition may still play a major role. Limited dispersal may favour altruism because it increases relatedness between

altruistic individuals. But it also increases relatedness between competitors, which opposes the selection for altruism (see, e.g. Taylor 1992; Wilson et al. 1992).

Both species in our experiment had fewer individuals in groups of relatives compared to groups of non-relatives suggesting that groups of related individuals suffered increased mortality compared to groups of non-related individuals. These results further suggested negative sibling competition effects, supporting the resource-partitioning hypothesis and are in line with previous studies on sibling competition in plants (Willson et al. 1987; Kelley 1989; Argyres and Schmitt 1992; Karron and Marshall 1993). Within groups of related individuals, the strong competitor produced on average the highest individual biomass (vegetative and reproductive). However, because mortality was highest in intrafamily aggregations, and because there was no difference in the overall biomass production between groups of relatives and groups of non-relatives, we interpret this result as a simple density dependent effect rather than with kin selection.

Finally, alternative explanations for the relatively weak effects of spatial patterns at the level of seed families and different responses of the species to intrafamily aggregation must be considered. For example, there may be little or no genetic variation in our populations. Because we do not have any knowledge about the genetic variation in the different species we cannot really assess this possible explanation. Moreover, differences in seed weight of the two species do certainly not provide the only explanation for different responses to intrafamily aggregation. The different mating systems among the target species (i.e. highly selfing for *Capsella* and predominately outcrossing for *Stachys*) might provide an alternative explanation for some of the different responses of the species to intrafamily aggregation. That is, selfed seeds of *Capsella* are genetically more similar than outcrossed seeds of *Stachys*. Therefore, related individuals of *Capsella* might have suffered more intense sibling competition compared to *Stachys*, resulting in less resource partitioning and consequently reduced biomass production in intrafamily aggregations.

In summary, we provided further evidence that weak competitors increased their fitness within neighbourhoods of conspecifics compared to neighbourhoods of heterospecifics. In contrast, intraspecific aggregation decreased the performance of strong competitors. An additional aggregation at the level of seed families produced less clear results. Nevertheless, groups of relatives tended to perform worse than groups of non-relatives in *Capsella* whereas in *Stachys*, groups of relatives tended to perform slightly better than groups of non-relatives. Therefore our findings tended to support the resource partitioning hypothesis, rather than the kin selection hypothesis. However, there are many other factors (e.g. habitat selection) that may affect the way individuals interact and aggregate, and the circumstances under which neighbourhood competition between related individuals may lead to kin selection in plants remain largely unexplored. Thus, the consequences of genetic substructuring for species coexistence and its ecological and evolutionary implication in plant population dynamics merits further investigation.

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Appendix

Appendix 1 Results of ANOVA for *Capsella bursa-pastoris* (top) and *Stachys annua* (bottom) using all plots

Source of variation	df	Total	Average								
			Number of individuals		Vegetative biomass		Reproductive biomass				
			F	P	F	P	F	P			
<i>Capsella bursa-pastoris</i>											
Block	1	1.4	0.271	0.1	0.760	0.2	0.667	6.0	0.039	13.0	0.007
Spatial pattern	2	1.7	0.236	3.6	0.078	10.0	0.007	13.6	0.003	36.2	<0.001
Random versus aggregated ^a	1	1.7	0.223	6.3	0.037	19.6	0.002	27.0	<0.001	72.3	<0.001
Intraspecific versus intrafamily aggregated ^a	1	1.7	0.225	0.9	0.380	0.3	0.580	0.3	0.616	0.2	0.694
Residual	8										
Total	11										
<i>Stachys annua</i>											
Block	1	6.2	0.038	0.7	0.427	0.0	0.998	6.2	0.038	6.1	0.039
Spatial pattern	2	0.7	0.510	0.1	0.878	0.2	0.842	0.3	0.721	0.3	0.721
Random versus aggregated ^a	1	0.7	0.415	0.0	0.899	0.1	0.833	0.1	0.778	0.1	0.778
Intraspecific versus intrafamily aggregated ^a	1	0.7	0.418	0.3	0.632	0.3	0.596	0.6	0.462	0.6	0.462
Residual	8										
Total	11										

Effects of spatial patterns (random versus two levels of aggregation) on total (sum over plots) number of individuals, total and average (per individual) aboveground vegetative and reproductive biomass production were tested against plot-level residuals

df degrees of freedom, F F-value (variance ratio), P error probability

^a Linear contrasts to separate the effects of spatial pattern into differences between random versus aggregated (i.e. intraspecific and intrafamily aggregation) and intraspecific versus intrafamily aggregation effects

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